

SEQUENCE LISTING

Tudan, Christopher R. Merzouk, Ahmed Arab, Lakhdar Saxena, Geeta Eaves, Connie J. Cashman, Johanne Clark-Lewis, Ian Salari, Hassan University of British Columbia Chemokine Therapeutics Corporation <120> CXC Chemokine Receptor 4 Agonist Peptides

- <130> 080421-000100US
- <140> US 10/086,177
- <141> 2002-02-26
- <150> CA 2,305,036
- <151> 2000-04-12
- <150> US 60/232,425
- <151> 2000-09-14
- <150> CA 2,335,109
- <151> 2001-02-23
- <150> US 09/835,107
- <151> 2001-04-12
- <160> 214
- <170> PatentIn version 3.3
- <210> 1
- <211> 67
- <212> PRT
- <213> Homo sapiens
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- Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Glu Ser 1
- His Val Ala Arg Ala Asn Val Lys His Leu Lys Ile Leu Asn Thr Pro 20
- Asn Cys Ala Leu Gln Ile Val Ala Arg Leu Lys Asn Asn Asn Arg Gln 40
- Val Cys Ile Asp Pro Lys Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys 50
- Ala Leu Asn
- 65
- <210> 2
- <211> 93
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- <213> Homo sapiens

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Cys Leu Ser Asp Gly Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys
                               25
Arg Phe Phe Glu Ser His Val Ala Arg Ala Asn Val Lys His Leu Lys
                           40
                                               45
Ile Leu Asn Thr Pro Asn Cys Ala Leu Gln Ile Val Ala Arg Leu Lys
                                           60
                       55
Asn Asn Arg Gln Val Cys Ile Asp Pro Lys Leu Lys Trp Ile Gln
                                       75
                   70
Glu Tyr Leu Glu Lys Ala Leu Asn Lys Arg Phe Lys Met
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Cys Leu Ser Asp Gly Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys
                               25
Arg Phe Phe Glu Ser His Val Ala Arg Ala Asn Val Lys His Leu Lys
                           40
Ile Leu Asn Thr Pro Asn Cys Ala Leu Gln Ile Val Ala Arg Leu Lys
                       55
Asn Asn Asn Arg Gln Val Cys Ile Asp Pro Lys Leu Lys Trp Ile Gln
                   70
                                       75
65
Glu Tyr Leu Glu Lys Ala Leu Asn Lys Arg Phe Lys Met
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Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Glu Ser
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His
<210> 5
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      6
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<220>
<223> CXCR4 agonist sequence motif within 20 amino acids
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of the N-terminus

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<400> 5
Arg Phe Phe Glu Ser His
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      synthetic SDF-1 peptide analogue CXCR4 agonist
<223>
<400> 6
Lys Pro Val Ser Leu Ser Tyr Arg Cys
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       CTCE9901
<220>
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       residues in position 7 of two SEQ ID NO:7 peptides
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Lys Pro Val Ser Leu Ser Tyr Arg Cys
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       (Compound #3)
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<223> Xaa = Lys whose epsilon amino group forms a covalent amide
       bond with the alpha amino group of Cys at position 9 of
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Lys Pro Val Ser Leu Ser Tyr Arg Cys Xaa
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<223>
       (Compound #3)
<220>
<221> MOD RES
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      (9)..(9)
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      bond with the epsilon amino group of Lys at position 10
      of KPVSLSYRCX (SEQ ID NO:8), thereby forming a dimer
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Lys Pro Val Ser Leu Ser Tyr Arg Xaa
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      acids 1-8
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<221> MOD RES
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      (9)..(9)
      Xaa = Lys whose epsilon amino group forms a covalent amide
<223>
       bond with the alpha amino group of Arg at position 8 of
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Lys Pro Val Ser Leu Ser Tyr Arg Xaa
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      (8)..(8)
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       bond with the epsilon amino group of Lys at position 9
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Lys Pro Val Ser Leu Ser Tyr Xaa
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            20
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<223> Gly in positions 17 and/or 18 may independently be
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                                25
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       amide
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<221> MISC FEATURE
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<223> Gly in position 17 may be present or absent
<220>
<221> MOD RES
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<223> AMIDATION
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Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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      amide, CTCE0017, Compound A
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<223> Gly in positions 17 and/or 18 may independently be
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                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
<210> 16
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<220>
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His Gly Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu
            20
                                25
Asn
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      34
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<223>
       acid
<220>
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<223> Gly in positions 20 and/or 21 may independently be
       present or absent
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Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Glu Ser
His Gly Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala
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Leu Asn
<210> 18
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<221> MOD RES
<222> (33)..(33)
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1
His Gly Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu
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Asn
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       present or absent
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<221> MOD RES
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Leu Asn
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                                25
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      (20)..(24)
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
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       CTCE0022
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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       CTCE0021, Compound B
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      (20)..(24)
      side chain cyclized using lactam formation
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       MOD RES
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      (31)..(31)
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                                    10
Gly Gly Leu Lys Trp Ile Gln Asp Tyr Leu Glu Lys Ala Leu Asn
                                25
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<223>
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       amide
<220>
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      side chain cyclized using lactam formation
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      MOD RES
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      (31) .. (31)
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Gly Gly Leu Lys Trp Ile Gln Asp Tyr Leu Glu Lys Ala Leu Asn
            20
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      present or absent
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                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
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       SDF-1(1-14)-(G)-4-SDF-1(55-67)-C9/C11-cyclic amide
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<223> Gly in positions 17 and/or 18 may be independently
       present or absent
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       MOD RES
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       (31)..(31)
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<223> AMIDATION
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                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
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      present or absent
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Gly Gly Ser Lys Pro Gly Val Ile Phe Leu Thr Lys Arg Ser Arg Gln
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                                                    30
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Val
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       present or absent
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                                    10
Gly Gly Cys Cys Phe Ser Tyr Thr Ser Arg Gln Ile Pro Gln Asn Phe
                                25
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Ile Ala Asp Tyr Phe Glu Thr Ser Ser Gln Cys Ser Lys Pro Gly Val
                                                45
Ile Phe Leu Thr Lys Arg Ser Arg Gln Val
                        55
    50
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                                    10
                5
Gly Gly Glu Glu Trp Val Gln Lys Tyr Val Asp Asp Leu Glu Leu Ser
                                25
            20
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Ala

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       Xaa = lysinamide whose epsilon amino group forms a
<223>
       covalent amide bond with the alpha amino group of Arg at
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       dimer
Lys Pro Val Ser Leu Ser Tyr Arg Xaa
                5
<210>
       32
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<223>
       bond with the epsilon amino group of lysinamide at
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<400> 32
Lys Pro Val Ser Leu Ser Tyr Xaa
<210>
       33
<400>
000
<210> 34
<400> 34
000
<210>
       35
       31
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       synthetic CXCR4 agonist SDF-1-derived cyclic amide
<223>
       (E24/K28)
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<223> Xaa = an amino acid that may be either an L-Pro or a
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                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
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       (K20/E24)
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      Lys may be modified with a substituent that may be a
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      (2)..(2)
<223> Xaa = an amino acid that may be either an L-Pro or a
       D-Pro moiety
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      D-Leu Moiety
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                                25
            20
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       acid
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       (24)..(28)
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Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Gly Gly
                                    10
1
                5
Gly Gly Leu Lys Trp Ile Gln Asp Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
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Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Asp Tyr Leu Glu Lys Ala Leu Asn
                                25
<210> 39
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       acid
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Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Gly Gly
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Gly Gly Leu Xaa Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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Sec. 1

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Gly Gly Leu Xaa Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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       amide
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Xaa Ala Leu Asn
                                25
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<223>
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       & E24->D)-cyclic acid
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                                    10
Gly Gly Leu Xaa Trp Ile Gln Asp Tyr Leu Glu Lys Ala Leu Asn
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       & E24->D)-cyclic amide
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<223> AMIDATION
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Gly Gly Leu Xaa Trp Ile Gln Asp Tyr Leu Glu Lys Ala Leu Asn
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Gly Gly Leu Lys Trp Ile Gln Asp Tyr Leu Glu Xaa Ala Leu Asn
                                25
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       & E24->D)-cyclic amide
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<223> Xaa = Orn
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Gly Gly Leu Lys Trp Ile Gln Asp Tyr Leu Glu Xaa Ala Leu Asn
            20
<210> 47
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<212> PRT
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1
Gly Gly Ser Lys Pro Gly Val Ile Phe Leu Thr Lys Arg Ser Arg Gln
                                25
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Val
<210> 48
<211> 58
<212> PRT
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<222>
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Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Gly Gly
                                    10
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                5
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Gly Gly Cys Cys Phe Ser Tyr Thr Ser Arg Gln Ile Pro Gln Asn Phe
                                25
Ile Ala Asp Tyr Phe Glu Thr Ser Ser Gln Cys Ser Lys Pro Gly Val
                            40
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Ile Phe Leu Thr Lys Arg Ser Arg Gln Val
                        55
<210> 49
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<212> PRT
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<220>
<221> MOD RES
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Gly Gly Glu Glu Trp Val Gln Lys Tyr Val Asp Asp Leu Glu Leu Ser
                                25
Ala
<210> 50
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<223> Xaa = an amino acid that may be either an L-Leu or a
       D-Leu moiety
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<220>
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<222> (24)..(28)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
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<222>
<223> AMIDATION
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                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 51
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<223>
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<220>
<221> MISC FEATURE
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<220>
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      (2)..(2)
<223> Xaa = an amino acid that may be either an L-Pro or a
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      Xaa = an amino acid that may be either an L-Leu or a
       D-Leu moiety
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<221>
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<223>
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       as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 51
Lys Xaa Val Ser Xaa Ser Tyr Arg Cys Pro Cys Arg Phe Xaa
1
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<210> 52
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<212> PRT
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<220>
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<223>
      E24/K28-cyclic amide
<220>
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<222>
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      Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
<223>
       (SEQ ID NO:51) via a moiety providing covalent attachment
      between N and C terminal portions of the peptides, such
      as NH-2-(CH-2)-n-COOH (n = 0-20)
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<222> (6)..(10)
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<222> (13)..(13)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                5
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<210> 53
<211> 31
<212> PRT
<213> Artificial Sequence
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<223> synthetic CXCR4 agonist SDF-1-derived K20/E24-cyclic
       amide
<220>
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      hydrogen, alkyl, aryl or polyethylene glycol (PEG) moiety
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      (2)..(2)
      Xaa = an amino acid that may be either an L-Pro or a
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       (5)..(5)
      Xaa = an amino acid that may be either an L-Leu or a
       D-Leu moiety
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Lys Xaa Val Ser Xaa Ser Tyr Arg Cys Pro Cys Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 54
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<223>
       K20/E24-cyclic amide
<220>
<221> MISC_FEATURE
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<223> Lys may be modified with a substituent that may be a
       hydrogen, alkyl, aryl or polyethylene glycol (PEG) moiety
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<223>
       D-Pro moiety
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      Xaa = an amino acid that may be either an L-Leu or a
<223>
       D-Leu moiety
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<223>
       (SEQ ID NO:55) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 54
Lys Xaa Val Ser Xaa Ser Tyr Arg Cys Pro Cys Arg Phe Xaa
                5
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<210> 55
<211> 13
<212> PRT
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<223>
       K20/E24-cyclic amide
<220>
<221>
      MOD RES
<222>
      (1)..(1)
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       (SEQ ID NO:54) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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<223> AMIDATION
<400> 55
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                5
<210> 56
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      side chain cyclized using lactam formation
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<223> AMIDATION
<400> 56
Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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<212> PRT
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Phe Arg Phe Phe Gly Gly
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
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       as NH-2-(CH-2)-n-COOH (n = 0-20)
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Phe Arg Phe Xaa
                                    10
                5
<210> 59
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<212>
      PRT
      Artificial Sequence
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      MOD RES
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       (SEQ ID NO:58) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 60
<211> 31
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<223> side chain cyclized using lactam formation
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Phe Arg Phe Phe Gly Gly
                5
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
<210> 61
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      (14)..(14)
<222>
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       (SEQ ID NO:62) via a moiety providing covalent attachment
      between N and C terminal portions of the peptides, such
      as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 61
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Phe Arg Phe Xaa
<210> 62
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<220>
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       (SEQ ID NO:61) via a moiety providing covalent attachment
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      as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 63
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<223>
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      side chain cyclized using lactam formation
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<221> MOD RES
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      (31)..(31)
<223> AMIDATION
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<400> 63
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro His Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
<210> 64
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<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:65) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro His Arg Phe Xaa
                5
                                    10
<210> 65
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                    10
                5
<210> 66
      31
<211>
<212> PRT
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<223>
<220>
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                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
<210> 67
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<223>
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       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro His Arg Phe Phe
                5
<210> 68
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<223> portion of synthetic CTCE0021-like analog CXCR4 agonist
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<222>
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      as NH-2-(CH-2)-n-COOH (n = 0-20)
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<400> 68
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
               5
<210> 69
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<223> AMIDATION
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Trp Arg Phe Phe Gly Gly
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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<210> 70
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<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:71) via a moiety providing covalent attachment
      between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 70
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Trp Arg Phe Xaa
<210> 71
<211> 13
<212> PRT
<213> Artificial Sequence
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<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:70) via a moiety providing covalent attachment
      between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 72
<211> 31
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<223> side chain cyclized using lactam formation
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<223> AMIDATION
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Trp Arg Phe Phe Gly Gly
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
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       (SEQ ID NO:74) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 73
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Trp Arg Phe Xaa
                5
<210> 74
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<220>
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<223>
       (SEQ ID NO:73) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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      side chain cyclized using lactam formation
<220>
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<222> (13)..(13)
<223> AMIDATION
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<400> 74
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 75
<211> 31
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<400> 75
Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Ala Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
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<221> MOD RES
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<222>
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:77) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Ala Arg Phe Xaa
                                    10
                5
<210> 77
<211>
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<212>
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       (SEQ ID NO:76) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 78
<211> 31
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<223> AMIDATION
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Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Ala Arg Phe Phe Gly Gly
               5
1
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
<210> 79
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       (SEQ ID NO:80) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Ala Arg Phe Xaa
                5
<210> 80
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       (SEQ ID NO:79) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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      AMIDATION
<223>
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                5
                                    10
<210> 81
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<400> 81
Lys Pro Val Ser Leu Ser Tyr Arg His Pro Ala Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
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<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:83) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 82
Lys Pro Val Ser Leu Ser Tyr Arg His Pro Ala Arg Phe Xaa
                5
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<210> 83
<211> 13
<212> PRT
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      Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
<223>
       (SEQ ID NO:82) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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<223>
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<223> AMIDATION
<400> 83
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                    10
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<210> 84
<211> 31
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<221> MOD RES
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<223> AMIDATION
<400> 84
Lys Pro Val Ser Leu Ser Tyr Arg His Pro Ala Arg Phe Phe Gly Gly
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
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      Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:86) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
Lys Pro Val Ser Leu Ser Tyr Arg His Pro Ala Arg Phe Xaa
                                    10
<210>
      86
<211>
      13
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      PRT
      Artificial Sequence
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      (SEQ ID NO:85) via a moiety providing covalent attachment
      between N and C terminal portions of the peptides, such
      as NH-2-(CH-2)-n-COOH (n = 0-20)
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<221> MISC FEATURE
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<221> MOD RES
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      (13)..(13)
<223> AMIDATION
<400> 86
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 87
<211> 31
<212> PRT
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<223> side chain cyclized using lactam formation
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<221> MOD RES
      (31)..(31)
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<223> AMIDATION
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Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Ala Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 88
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       (SEQ ID NO:89) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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<400> 88
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Ala Arg Phe Xaa
<210> 89
<211> 13
<212> PRT
<213> Artificial Sequence
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       (SEQ ID NO:88) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 90
<211> 31
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Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Ala Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 91
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       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 91
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Ala Arg Phe Xaa
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<210> 92
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       as NH-2-(CH-2)-n-COOH (n = 0-20)
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<223> AMIDATION
<400> 92
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                    10
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<210> 93
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<223> AMIDATION
<400> 93
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Tyr Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
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<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:95) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
      as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 94
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Tyr Arg Phe Xaa
                                    10
<210> 95
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      13
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       (SEQ ID NO:94) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
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      (13)..(13)
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<400> 95
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 96
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                                    10
1
                5
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
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<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
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      between N and C terminal portions of the peptides, such
      as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 97
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Tyr Arg Phe Xaa
<210> 98
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<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:97) via a moiety providing covalent attachment
      between N and C terminal portions of the peptides, such
      as NH-2-(CH-2)-n-COOH (n = 0-20)
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<400> 98
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 99
      31
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      side chain cyclized using lactam formation
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<221> MOD RES
      (31)..(31)
<222>
<223> AMIDATION
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<400> 99
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Tyr Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 100
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<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:101) via a moiety providing covalent attachment
       between N and C terminal portions of the peptides, such
       as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 100
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Tyr Arg Phe Xaa
                5
                                    10
<210> 101
<211> 13
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       (SEQ ID NO:100) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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                5
                                    10
<210> 102
<211> 31
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<223> Lys is modified with polyethylene glycol (PEG)
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                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
<210> 103
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      (14)..(14)
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       (SEQ ID NO:104) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 103
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Tyr Arg Phe Xaa
                5
                                    10
<210> 104
      13
<211>
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<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:103) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<221> MISC_FEATURE
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Ala Arg Phe Phe Gly Gly
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
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<210> 106
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      Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
<223>
       (SEQ ID NO:107) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Ala Arg Phe Xaa
<210> 107
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<223> AMIDATION
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Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Ala Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
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<221> MOD RES
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<223>
       (SEQ ID NO:110) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Ala Arg Phe Xaa
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                                    10
1
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<223>
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       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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                                    10
<210> 111
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Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Phe Arg Phe Phe Gly Gly
               5
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
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       (SEQ ID NO:113) via a moiety providing covalent
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Phe Arg Phe Xaa
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                5
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       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<222>
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<223> side chain cyclized using lactam formation
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Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Phe Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
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       (SEQ ID NO:116) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<400> 115
Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Phe Arg Phe Xaa
<210> 116
<211> 13
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      (SEQ ID NO:115) via a moiety providing covalent
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 117
<211> 31
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<223> AMIDATION
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Lys Pro Val Ser Leu Ser Tyr Arg His Pro His Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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<210> 118
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      (SEQ ID NO:119) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
Lys Pro Val Ser Leu Ser Tyr Arg His Pro His Arg Phe Xaa
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       (SEQ ID NO:118) via a moiety providing covalent
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 120
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Lys Pro Val Ser Leu Ser Tyr Arg His Pro His Arg Phe Phe Gly Gly
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
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<220>
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<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:122) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
Lys Pro Val Ser Leu Ser Tyr Arg His Pro His Arg Phe Xaa
                5
                                    10
<210> 122
<211>
      13
      PRT
<212>
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      (1)..(1)
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       (SEQ ID NO:121) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 123
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Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Trp Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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<210> 124
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       (SEQ ID NO:125) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Trp Arg Phe Xaa
                5
                                    10
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<210> 125
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      (SEQ ID NO:124) via a moiety providing covalent
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 126
<211> 31
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Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Trp Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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      (SEQ ID NO:128) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Trp Arg Phe Xaa
                5
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<400> 128
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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<210> 129
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                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
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       (SEQ ID NO:131) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Xaa
                5
<210> 131
      13
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       (SEQ ID NO:130) via a moiety providing covalent
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<210> 132
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                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
           20
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      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Phe Arg Phe Xaa
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       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<210> 135
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Phe Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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<210> 136
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Phe Arg Phe Xaa
                5
<210> 137
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       (SEQ ID NO:136) via a moiety providing covalent
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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      (13)..(13)
<223>
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro His Arg Phe Xaa
                                    10
               5
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro His Arg Phe Phe Gly Gly
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 142
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro His Arg Phe Xaa
<210> 143
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       (SEQ ID NO:142) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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      side chain cyclized using lactam formation
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<222>
<223> AMIDATION
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Trp Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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       (SEO ID NO:146) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 145
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Trp Arg Phe Xaa
<210> 146
<211> 13
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       (SEQ ID NO:145) via a moiety providing covalent
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 147
<211> 31
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Trp Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
<210> 148
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       (SEQ ID NO:149) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 148
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Trp Arg Phe Xaa
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<210> 149
<211>
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<212>
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       (SEQ ID NO:148) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 150
<211> 31
<212> PRT
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<223> AMIDATION
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Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Ala Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 151
<211>
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       Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
<223>
       (SEQ ID NO:152) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<400> 151
Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Ala Arg Phe Xaa
<210> 152
<211> 13
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       (SEQ ID NO:151) via a moiety providing covalent
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      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 153
<211> 31
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<223> AMIDATION
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<400> 153
Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Ala Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
<210> 154
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       (SEQ ID NO:155) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 154
Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Ala Arg Phe Xaa
                5
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<210> 155
<211> 13
<212> PRT
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<223>
       (SEQ ID NO:154) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
1
                5
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<210> 156
<211>
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<400> 156
Lys Pro Val Ser Leu Ser Tyr Arg His Pro Ala Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 157
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<222>
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:158) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 157
Lys Pro Val Ser Leu Ser Tyr Arg His Pro Ala Arg Phe Xaa
<210> 158
<211> 13
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      Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:157) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
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Lys Pro Val Ser Leu Ser Tyr Arg His Pro Ala Arg Phe Phe Gly Gly
1
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
<210> 160
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<222>
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:161) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 160
Lys Pro Val Ser Leu Ser Tyr Arg His Pro Ala Arg Phe Xaa
<210> 161
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<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:160) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 162
<211> 31
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      (31)..(31)
<223> AMIDATION
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<400> 162
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Ala Arg Phe Phe Gly Gly
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Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
<210> 163
<211> 14
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<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:164) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Ala Arg Phe Xaa
<210>
      164
<211>
      13
<212> PRT
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       (SEQ ID NO:163) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<220>
      MOD RES
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      (13)..(13)
<223> AMIDATION
<400> 164
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                5
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<210>
      165
<211>
      31
<212>
      PRT
<213> Artificial Sequence
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<220>
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<220>
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<400> 165
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Ala Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 166
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<212> PRT
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<221> MOD RES
<222>
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<223>
       (SEQ ID NO:167) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 166
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Ala Arg Phe Xaa
                5
                                    10
<210> 167
<211>
      13
<212>
      PRT
<213> Artificial Sequence
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<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
      (SEQ ID NO:166) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
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<222>
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<400> 167
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 168
<211> 31
<212> PRT
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<223>
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      side chain cyclized using lactam formation
<223>
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      (31)..(31)
<222>
<223> AMIDATION
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Tyr Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 169
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       (SEQ ID NO:170) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<400> 169
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Tyr Arg Phe Xaa
<210> 170
<211> 13
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<213> Artificial Sequence
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       (SEQ ID NO:169) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
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<223> side chain cyclized using lactam formation
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<222> (13)..(13)
<223> AMIDATION
<400> 170
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 171
<211> 31
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<223>
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<223> side chain cyclized using lactam formation
<220>
<221> MOD_RES
<222> (31)..(31)
<223> AMIDATION
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Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Tyr Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 172
<211>
      14
<212> PRT
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<223>
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<220>
<221> MOD RES
<222>
      (14)..(14)
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:173) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 172
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Tyr Arg Phe Xaa
                5
<210> 173
<211> 13
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<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
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      Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
<223>
       (SEQ ID NO:172) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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      side chain cyclized using lactam formation
<220>
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<223>
      AMIDATION
<400> 173
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                    10
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<210> 174
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<223> AMIDATION
<400> 174
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Tyr Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 175
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      (14)..(14)
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:176) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 175
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Tyr Arg Phe Xaa
<210> 176
<211> 13
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<213> Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
      MOD RES
<221>
<222>
       (1)..(1)
      Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:175) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<220>
<221> MISC_FEATURE
<222>
      (6)..(10)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
<222>
      (13)..(13)
<223> AMIDATION
<400> 176
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 177
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD RES
<222> (1)..(1)
<223> Lys is modified with polyethylene glycol (PEG)
<220>
<221> MISC FEATURE
<222> (24)..(28)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
      (31)...(31)
<222>
<223> AMIDATION
<400> 177
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Tyr Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
<210> 178
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD_RES
<222>
      (1)..(1)
<223> Lys is modified with polyethylene glycol (PEG)
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<220>
<221> MOD_RES
<222>
      (14)..(14)
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:179) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 178
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Tyr Arg Phe Xaa
<210> 179
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223>
      portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD_RES
<222>
      (1)..(1)
<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:178) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
<221> MISC FEATURE
<222> (6)..(10)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
<222> (13)..(13)
<223> AMIDATION
<400> 179
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 180
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
      synthetic CTCE0022-like analog CXCR4 agonist
<223>
<220>
<221> MISC FEATURE
<222>
      (24)..(28)
      side chain cyclized using lactam formation
<220>
<221> MOD RES
<222>
      (31)..(31)
<223> AMIDATION
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<400> 180
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Ala Arg Phe Phe Gly Gly
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 181
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD RES
<222> (14)..(14)
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
      (SEQ ID NO:182) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Ala Arg Phe Xaa
<210> 182
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD RES
<222>
      (1)..(1)
<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:181) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
<221> MISC_FEATURE
<222>
      (6)..(10)
      side chain cyclized using lactam formation
<223>
<220>
<221> MOD RES
      (13)..(13)
<222>
<223> AMIDATION
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 183
<211> 31
<212> PRT
<213> Artificial Sequence
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<220>
<223>
      synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD_RES
<222>
      (1)..(1)
<223> Lys is modified with polyethylene glycol (PEG)
<220>
<221> MISC_FEATURE
<222> (24)..(28)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
      (31)..(31)
<222>
<223> AMIDATION
<400> 183
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Ala Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 184
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD RES
<222> (1)..(1)
<223> Lys is modified with polyethylene glycol (PEG)
<220>
<221> MOD_RES
      (14)..(14)
<222>
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:185) via a moiety providing covalent
      attachment between N and C terminal portions of the \,
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 184
Lys Pro Val Ser Leu Ser Tyr Arg Tyr Pro Ala Arg Phe Xaa
                5
                                    10
1
<210> -185
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
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<220>
<221> MOD_RES
<222>
      (1)..(1)
<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:184) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
<221> MISC_FEATURE
      (6)..(10)
<222>
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
<222>
      (13)..(13)
<223> AMIDATION
<400> 185
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 186
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
      synthetic CTCE0022-like analog CXCR4 agonist
<223>
<220>
<221> MISC FEATURE
<222> (24)..(28)
<223> side chain cyclized using lactam formation
<220>
<221>
      MOD RES
      (31)..(31)
<222>
<223> AMIDATION
<400> 186
Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Phe Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 187
<211> 14
<212> PRT
      Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221>
      MOD RES
<222>
      (14)..(14)
      Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:188) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
```

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<400> 187
Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Phe Arg Phe Xaa
<210> 188
<211>
      13
<212>
      PRT
<213> Artificial Sequence
<220>
      portion of synthetic CTCE0022-like analog CXCR4 agonist
<223>
<220>
<221> MOD_RES
<222>
      (1)..(1)
<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:187) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
<221> MISC_FEATURE
<222> (6)..(10)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
<222>
      (13)..(13)
<223> AMIDATION
<400> 188
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 189
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
      synthetic CTCE0022-like analog CXCR4 agonist
<223>
<220>
<221> MOD_RES
<222>
      (1)..(1)
<223> Lys is modified with polyethylene glycol (PEG)
<220>
      MISC_FEATURE
<221>
      (24)..(28)
<222>
      side chain cyclized using lactam formation
<223>
<220>
      MOD_RES
<221>
<222>
      (31)..(31)
<223>
      AMIDATION
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<400> 189
 Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Phe Arg Phe Phe Gly Gly
 Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                 25
 <210> 190
 <211> 14
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> portion of synthetic CTCE0022-like analog CXCR4 agonist
 <220>
 <221> MOD RES
 <222> (1)..(1)
 <223> Lys is modified with polyethylene glycol (PEG)
 <220>
 <221> MOD RES
       (14)..(14)
 <222>
 <223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
        (SEQ ID NO:191) via a moiety providing covalent
        attachment between N and C terminal portions of the
        peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
 <400> 190
 Lys Pro Val Ser Leu Ser Tyr Arg Phe Pro Phe Arg Phe Xaa
 1
                 5
 <210> 191
 <211>
       13
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> portion of synthetic CTCE0022-like analog CXCR4 agonist
· <220>
       MOD RES
 <221>
 <222>
       (1)..(1)
        Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
 <223>
        (SEQ ID NO:190) via a moiety providing covalent
        attachment between N and C terminal portions of the
        peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
 <220>
 <221>
       MISC FEATURE
 <222>
       (6)..(10)
        side chain cyclized using lactam formation
 <220>
       MOD RES
 <221>
        (13)..(13)
 <222>
       AMIDATION
 <223>
 <400> 191
 Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                     10
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<210> 192
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
      synthetic CTCE0022-like analog CXCR4 agonist
<223>
<220>
<221> MISC_FEATURE
<222> (24)..(28)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
<222> (31)..(31)
<223> AMIDATION
<400> 192
Lys Pro Val Ser Leu Ser Tyr Arg His Pro His Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
           20
<210> 193
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD RES
<222>
      (14)..(14)
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:194) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 193
Lys Pro Val Ser Leu Ser Tyr Arg His Pro His Arg Phe Xaa
                                    10
<210> 194
<211> 13
<212>
      PRT
<213> Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221>
      MOD_RES
<222>
      (1)..(1)
      Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:193) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<220>
<221> MISC_FEATURE
<222> (6)..(10)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
<222> (13)..(13)
<223> AMIDATION
<400> 194
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 195
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD RES
<222> (1)..(1)
<223> Lys is modified with polyethylene glycol (PEG)
<220>
<221> MISC_FEATURE
<222> (24)..(28)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
      (31)..(31)
<222>
<223> AMIDATION
<400> 195
Lys Pro Val Ser Leu Ser Tyr Arg His Pro His Arg Phe Phe Gly Gly
1
               5
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
            20
<210> 196
<211>
      14
<212>
      PRT
<213> Artificial Sequence
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD RES
<222>
      (1)..(1)
<223> Lys is modified with polyethylene glycol (PEG)
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<220>
<221> MOD_RES
<222>
      (14)..(14)
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:197) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 196
Lys Pro Val Ser Leu Ser Tyr Arg His Pro His Arg Phe Xaa
<210> 197
<211> 13
<212> PRT
<213> Artificial Sequence
      portion of synthetic CTCE0022-like analog CXCR4 agonist
<223>
<220>
<221> MOD RES
<222>
      (1)..(1)
<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:196) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
<221> MISC_FEATURE
<222>
      (6)..(10)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
      (13)..(13)
<222>
<223> AMIDATION
<400> 197
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                5
<210> 198
<211>
      31
<212>
<213>
      Artificial Sequence
<220>
      synthetic CTCE0022-like analog CXCR4 agonist
<223>
<220>
<221> MISC_FEATURE
<222>
      (24)..(28)
      side chain cyclized using lactam formation
<400> 198
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Trp Arg Phe Phe Gly Gly
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                25
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<210> 199
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
      portion of synthetic CTCE0022-like analog CXCR4 agonist
<223>
<220>
<221> MOD RES
<222>
      (14)..(14)
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:200) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 199
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Trp Arg Phe Xaa
<210> 200
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD_RES
<222>
      (1)..(1)
<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:199) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
<221> MISC_FEATURE
<222>
      (6)..(10)
      side chain cyclized using lactam formation
<223>
<220>
<221>
      MOD RES
<222>
      (13)..(13)
<223> AMIDATION
<400> 200
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 201
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic CTCE0022-like analog CXCR4 agonist
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<220>
<221> MOD_RES
<222> (1)..(1)
<223> Lys is modified with polyethylene glycol (PEG)
<220>
<221> MISC_FEATURE
<222> (24)..(28)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
<222> (31)..(31)
<223> AMIDATION
<400> 201
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Trp Arg Phe Phe Gly Gly
               5
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
            20
                                25
<210> 202
<211> 14
<212> PRT
<213> Artificial Sequence
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
<221> MOD RES
<222> (1)..(1)
<223> Lys is modified with polyethylene glycol (PEG)
<220>
<221> MOD RES
<222>
      (14)..(14)
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:203) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 202
Lys Pro Val Ser Leu Ser Tyr Arg Trp Pro Trp Arg Phe Xaa
                5
                                    10
<210> 203
<211>
      13
      PRT
<212>
      Artificial Sequence
<220>
<223> portion of synthetic CTCE0022-like analog CXCR4 agonist
<220>
      MOD RES
<221>
<222>
      (1)..(1)
      Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
<223>
       (SEQ ID NO:202) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
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<220>
<221> MISC FEATURE
<222>
      (6)..(10)
<223> side chain cyclized using lactam formation
<220>
<221> MOD_RES
<222>
      (13)..(13)
<223> AMIDATION
<400> 203
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 204
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
     portion of synthetic CXCR4 agonist SDF-1-derived cyclic
<223>
       amide (E24/K28)
<220>
<221> MISC FEATURE
<222>
      (1)..(1)
<223> Lys may be modified with a substituent that may be a
      hydrogen, alkyl, aryl or polyethylene glycol (PEG) moiety
<220>
<221> MISC FEATURE
<222> (2)..(2)
<223> Xaa = an amino acid that may be either an L-Pro or a
      D-Pro moiety
<220>
<221> MISC FEATURE
<222>
      (5)..(5)
<223> Xaa = an amino acid that may be either an L-Leu or a
       D-Leu moiety
<220>
      MOD RES
<221>
<222>
      (14)..(14)
      Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
<223>
       (SEQ ID NO:205) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 204
Lys Xaa Val Ser Xaa Ser Tyr Arg Cys Pro Cys Arg Phe Xaa
                                    10
1
                5
<210> 205
<211>
      13
      PRT
<212>
<213> Artificial Sequence
<223> portion of synthetic CXCR4 agonist SDF-1-derived cyclic
       amide (E24/K28)
```

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<220>
<221> MOD RES
<222>
      (1)..(1)
<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:204) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
<221> MISC_FEATURE
<222>
      (6)..(10)
<223>
      side chain cyclized using lactam formation
<220>
<221>
      MOD RES
<222>
      (13)..(13)
<223>
      AMIDATION
<400> 205
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
<210> 206
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
      portion of synthetic CXCR4 agonist SDF-1-derived cyclic
<223>
       acid (K20/E24)
<220>
<221> MISC FEATURE
<222> (1)..(1)
<223> Lys may be modified with a substituent that may be a
       hydrogen, alkyl, aryl or polyethylene glycol (PEG) moiety
<220>
<221> MISC FEATURE
<222>
      (2)..(2)
<223> Xaa = an amino acid that may be either an L-Pro or a
       D-Pro moiety
<220>
<221> MISC FEATURE
<222>
      (5)..(5)
<223> Xaa = an amino acid that may be either an L-Leu or a
       D-Leu moiety
<220>
      MOD_RES
<221>
      (14)..(14)
<222>
      Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
<223>
       (SEQ ID NO:207) via a moiety providing covalent
       attachment between N and C terminal portions of the
       peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 206
Lys Xaa Val Ser Xaa Ser Tyr Arg Cys Pro Cys Arg Phe Xaa
                                    10
1
                5
```

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<210> 207
<211>
      13
<212>
      PRT
<213> Artificial Sequence
<220>
      portion of synthetic CXCR4 agonist SDF-1-derived cyclic
<223>
       acid (K20/E24)
<220>
<221> MOD_RES
<222>
      (1)..(1)
<223> Xaa = Leu linked to Phe at position 14 of KXVSXSYRCPCRFX
       (SEQ ID NO:206) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
<221> MISC FEATURE
<222> (6)..(10)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
<222>
      (13)..(13)
<223> AMIDATION
<400> 207
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                    10
                5
<210> 208
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic CXCR4 agonist
<220>
<221> MISC_FEATURE
      (17)..(18)
<222>
<223> Gly in positions 17 and/or 18 may be independently
      present or absent
<220>
<221> MISC FEATURE
      (20)..(24)
<222>
<223> side chain cyclized using lactam formation
<220>
      MOD RES
<221>
<222>
      (31)..(31)
<223>
      AMIDATION
<400> 208
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Phe Arg Phe Phe Gly Gly
                5
                                    10
Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                              25
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<210> 209
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223>
      portion of synthetic CXCR4 agonist
<220>
<221> MOD_RES
<222>
      (14)..(14)
<223> Xaa = Phe linked to Leu at position 1 of XKWIQEYLEKALN
       (SEQ ID NO:210) via a moiety providing covalent
       attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<400> 209
Lys Pro Val Ser Leu Ser Tyr Arg Ala Pro Phe Arg Phe Xaa
<210> 210
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> portion of synthetic CXCR4 agonist
<220>
<221> MOD RES
      (1)..(1)
<222>
<223> Xaa = Leu linked to Phe at position 14 of KPVSLSYRAPFRFX
       (SEQ ID NO:209) via a moiety providing covalent
      attachment between N and C terminal portions of the
      peptides, such as NH-2-(CH-2)-n-COOH (n = 0-20)
<220>
<221> MISC FEATURE
<222> (2)..(6)
<223> side chain cyclized using lactam formation
<220>
<221> MOD RES
      (13)..(13)
<222>
<223> AMIDATION
<400> 210
Xaa Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
                                    10
<210> 211
<211>
      4
<212>
      PRT
<213> Artificial Sequence
<220>
<223> 4 glycine linking moiety, [linker]
<400> 211
Gly Gly Gly Gly
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<210> 212
<211>
<212> PRT
<213> Artificial Sequence
<220>
<223> three or four glycine linker
<220>
<221> MISC_FEATURE
<222> (4)..(4)
<223> Gly in position 4 may be present or absent
<400> 212
Gly Gly Gly Gly
<210> 213
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> G-1-4 linker
<220>
<221> MISC FEATURE
<222> (2)..(4)
<223> Gly in positions 2-4 may be present or absent
<400> 213
Gly Gly Gly Gly
<210> 214
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> variable spacer monomer
<220>
<221> MISC_FEATURE
<222>
      (3)..(4)
<223> Gly in positions 3-4 may be present or absent
<400> 214
Gly Gly Gly Gly
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